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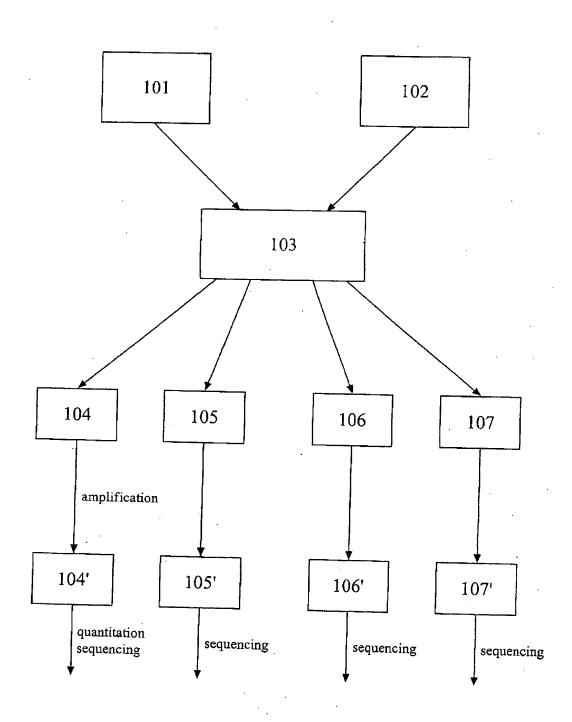


FIG. 1

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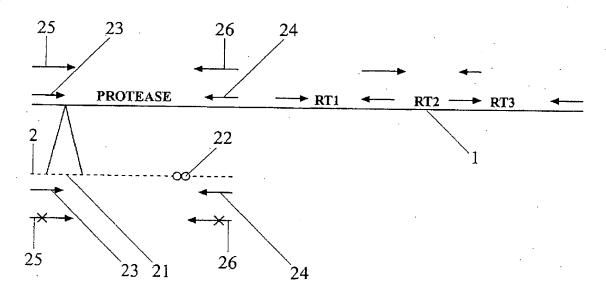


FIG. 2A

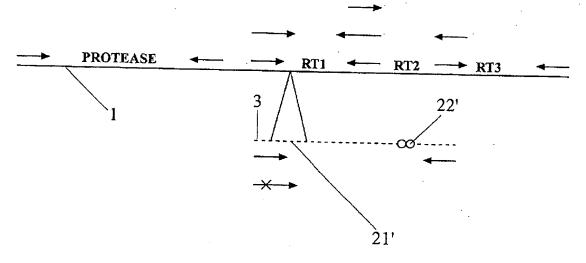


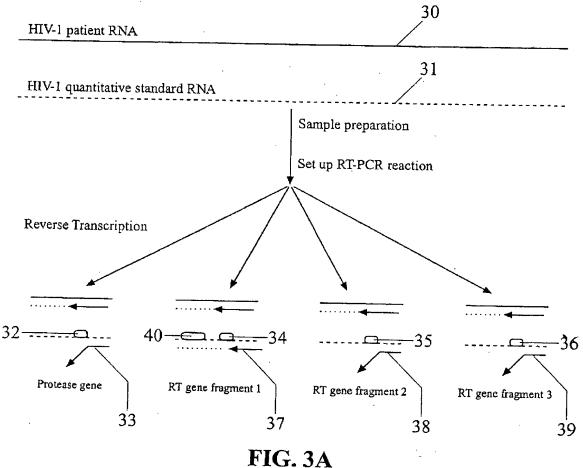
FIG. 2B

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Protease gene RT gene fragment 1 RT gene fragment 2 RT gene fragment 3 Polymerase Chain Reaction 

FIG. 3B



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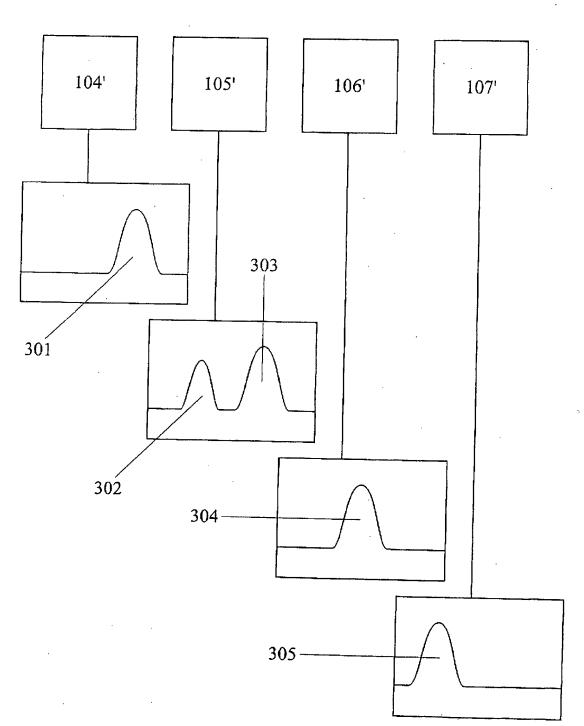


FIG. 3C

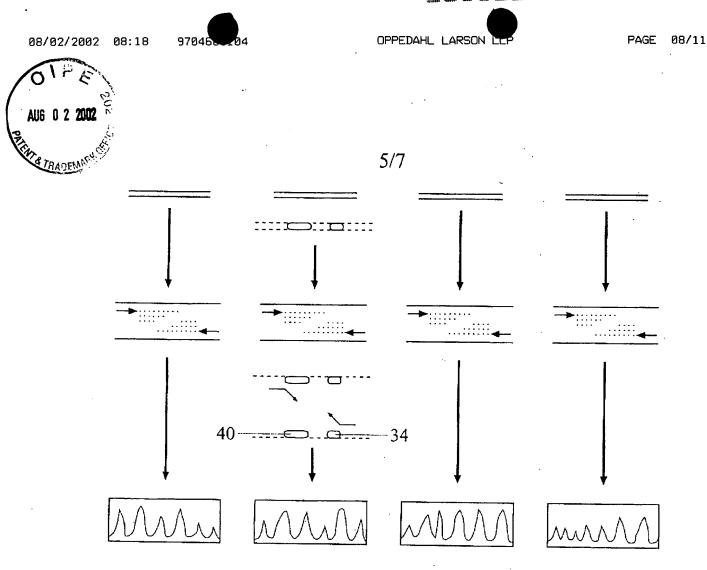
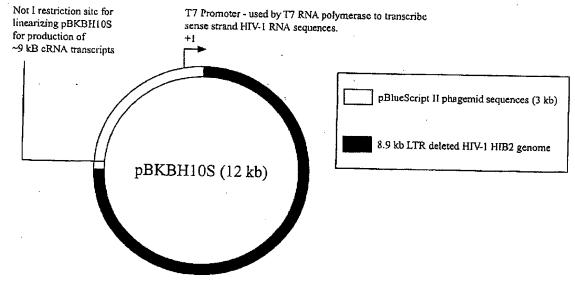


FIG. 3D



**FIG. 4** 

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5' to 3' direction of HIV-1 sense sequence RT562F-A primer (for sequencing) RT558F primer (for RT-PCR)

5' aaaagttaaacaatggccattgacag aagaaaaaataaaagcattagtagaaatttgtacagagatggaaaagga 3' base pairs 3' ttttcaatttgttaccggtaactgtc ttctttttatttcgtaatcatctttaaacatgtctctaccttttcct 5' 2606 to 2680 16 bp insertion added between nucleotides 2631 and 2632. tcctaaataaatagca The four terminal 3' nucleotides sequencing primer RT562F-A aggatttatttatcgt do not hybidize to sequences in the insert.

FIG. 5

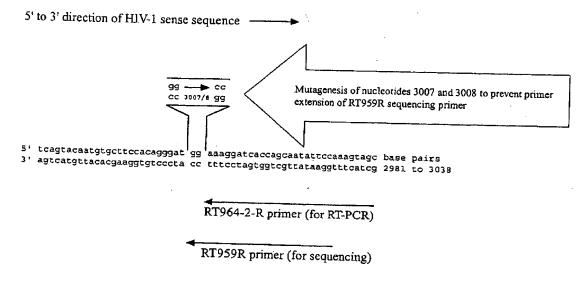
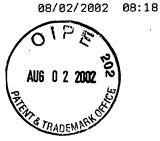


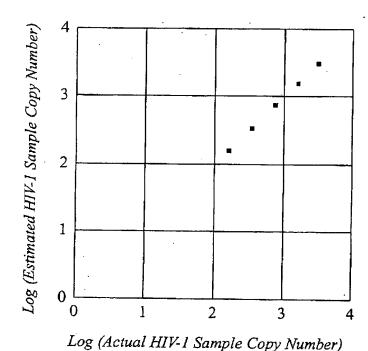
FIG. 6



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## Estimated vs. Actual HIV-1 Sample Copy Number



## Regression Statistics

slope: 1.060750

error of slope: 0.032348

y-Int.: -0.204487

error of y-Int.: 0.094912

x mean: 2.903090 y mean: 2.874965 x Std. Dev.: 0.475971 y Std. Dev.: 0.505588

R value: 0.998612

**FIG.** 7